

ONE DAY ONE GENOME



Isolated from a chronic wound

Staphylococcus saprophyticus

An emerging wound pathogen with biofilm and resistance traits

Quality of Genome Assembly and Annotation:
Results from indigenously developed **BHARAT** analysis pipeline: (**B**acterial **H**ybrid genome **A**ssembly and **R**apid **A**nnotation **T**oolset)

Table 1: Assembly Details	
Contigs	471
GC Content	32.65
Contig L50	109
Genome length	977,255 bp
Contig N50	2,342

Table 2: Annotated Genome Features	
CDS	1,454
tRNA	12
Repeat Regions	0
rRNA	2

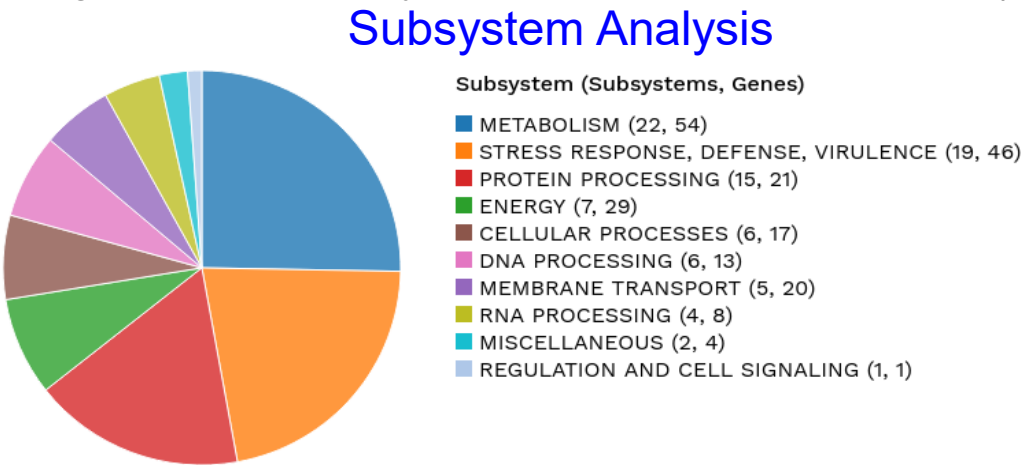


Table 3: Antimicrobial Resistance Genes	
AMR Mechanism	Genes
Antibiotic inactivation enzyme	FosB
Antibiotic resistance gene cluster,cassette,or operon	TcaR
Antibiotic target in susceptible species	Alr, Ddl, EF-G, folA, Dfr, folP, gyrA, gyrB, inhA, fabI, rpoB, rpoC, S12p
Protein altering cell wall charge conferring antibiotic resistance	GdpD
Efflux pump conferring antibiotic resistance	NorA, YkkCD
Regulator modulating expression of antibiotic resistance genes	LiaF, LiaR, LiaS

